

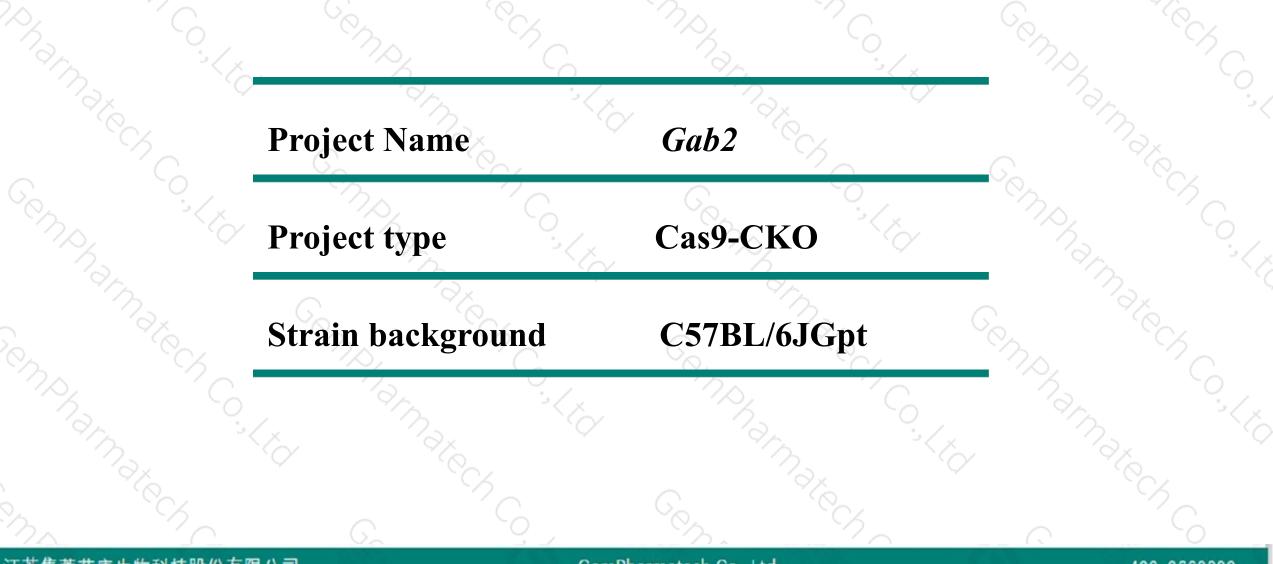
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Project Overview





江苏集萃药康生物科技股份有限公司

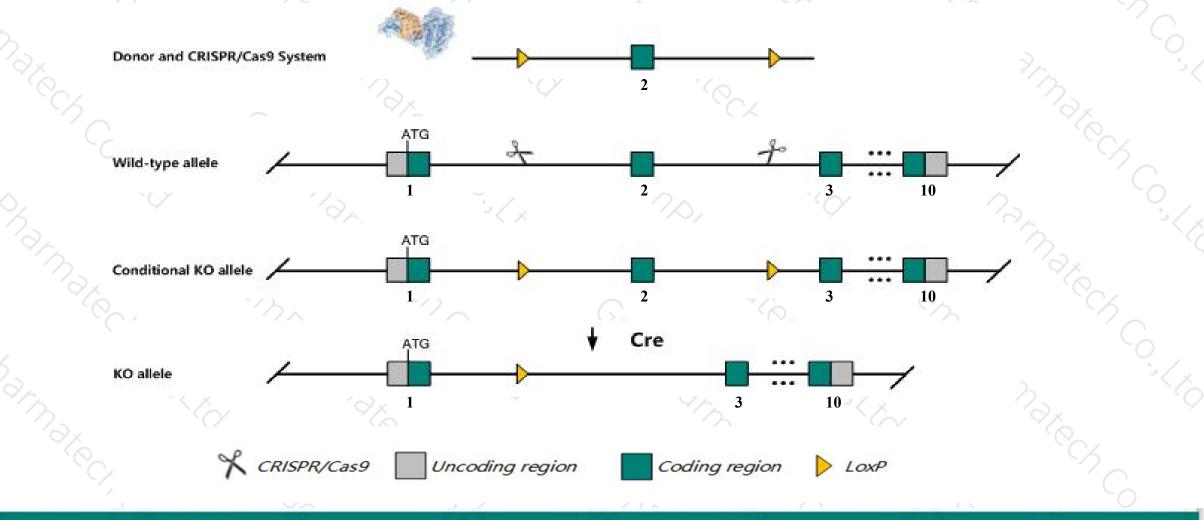
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Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Gab2 gene. The schematic diagram is as follows:



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 The Gab2 gene has 2 transcripts. According to the structure of Gab2 gene, exon2 of Gab2-201 (ENSMUST0000004622.6) transcript is recommended as the knockout region. The region contains 301bp coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Gab2* gene. The brief process is as follows:CRISPR/Cas9 system and Donor were microinjected into the fertilized eggs of C57BL/6JGpt mice.Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.

The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.



- According to the existing MGI data, Homozygotes for targeted null mutations exhibit impairments in passive cutaneous and systemic anaphylaxis, Fc gamma receptor-mediated phagocytosis, and mast cell development.
- The Gab2 gene is located on the Chr7. If the knockout mice are crossed with other mice strains to obtain double gene positive homozygous mouse offspring, please avoid the two genes on the same chromosome.
- This Strategy is designed based on genetic information in existing databases. Due to the complexity of biological processes, all risk of loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)



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Gab2 growth factor receptor bound protein 2-associated protein 2 [Mus musculus (house mouse)]

Gene ID: 14389, updated on 31-Jan-2019

Summary

Official Symbol	Gab2 provided by MGI
	growth factor receptor bound protein 2-associated protein 2 provided by MGI
Primary source	MGI:MGI:1333854
2.70	Ensembl:ENSMUSG0000004508
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;
	Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AI463667, D130058I17Rik, p97
Expression	Ubiquitous expression in ovary adult (RPKM 14.2), testis adult (RPKM 7.5) and 27 other tissues See more
Orthologs	human all



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The gene has 2 transcripts, all transcripts are shown below:

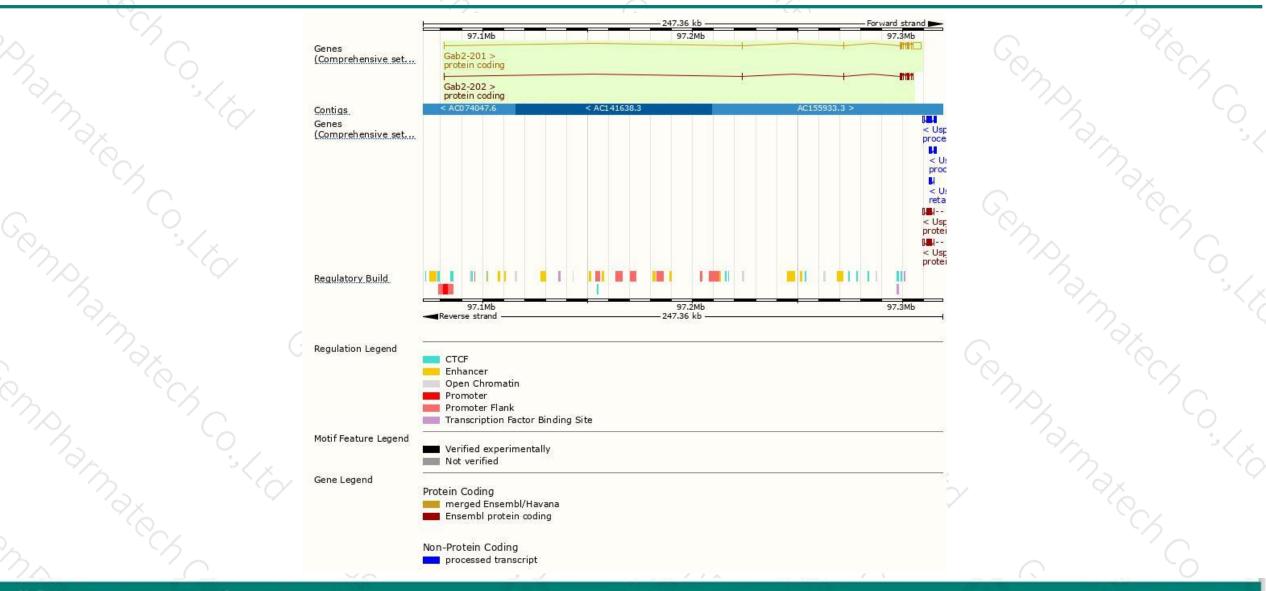
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Gab2-201	ENSMUST0000004622.6	6168	<u>666aa</u>	Protein coding	CCDS21455	Q3ZB57	TSL:1 GENCODE basic APPRIS P3
Gab2-202	ENSMUST00000206791.1	2065	<u>665aa</u>	Protein coding	CCDS85340	Q3ZB59	TSL:1 GENCODE basic APPRIS ALT2

The strategy is based on the design of *Gab2-201* transcript, The transcription is shown below

			- 227	.36 kb		For	ward strand
Gab2-201 >						L.	
Gab2-201 > protein coding							
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Genomic location distribution





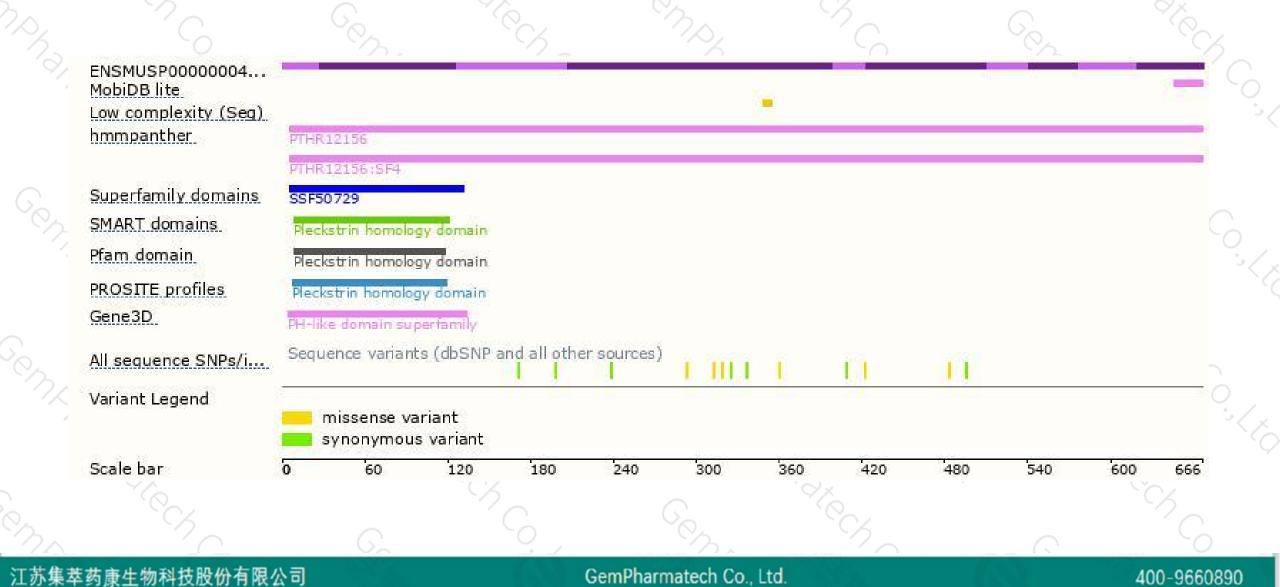
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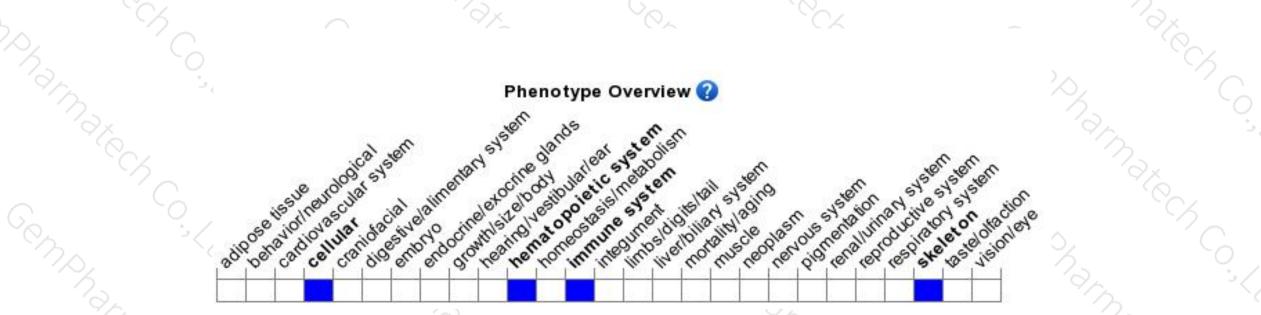
Protein domain





Mouse phenotype description(MGI)





Phenotypes affected by the gene are marked in blue.Data quoted from MGI database(http://www.informatics.jax.org/).

According to the existing MGI data, Homozygotes for targeted null mutations exhibit impairments in passive cutaneous and systemic anaphylaxis, Fc gamma receptor-mediated phagocytosis, and mast cell development.



If you have any questions, you are welcome to inquire. Tel: 400-9660890



